



1

SEQUENCE LISTING

<110> TSENG, RICHARD W.
SAMOSZUK, MICHAEL K.

<120> BCR-ABL GENE REARRANGMENT ASSAY METHOD

<130> 034827/0302

<140> 09/747,165

<141> 2000-12-22

<150> 60/173,050

<151> 1999-12-24

<160> 8

<170> PatentIn Ver. 2.1

<210> 1

<211> 19

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 1

cctcgagaa ctgcaaca

19

<210> 2

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 2

gagctgcaga tgctgaccaa

20

<210> 3

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer

<400> 3

tcagaccctg aggtctaaag tc

22

<210> 4

<211> 19

<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 4
gaaggtgaag gtcggagtc 19

<210> 5
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 5
gaagatggtg atgggatttc 20

<210> 6
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Probe

<400> 6
acacgacaac cgggcagtgc c 21

<210> 7
<211> 28
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Probe

<400> 7
tgctgtggac agtctggagt ttcacaca 28

<210> 8
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Probe

<400> 8
caagcttccc gttctcagcc 20